

IN THE CLAIMS

Claims 1-25 (Cancelled).

Claim 26 (Currently Amended): A process for producing L-amino acids comprising culturing a bacterial cell in a medium suitable for producing L-amino acids, wherein said bacterial cell comprises an attenuated lysR3 gene, wherein the lysR3 gene prior to being attenuated comprises SEQ ID NO:1, ~~SEQ ID NO:3~~, or a polynucleotide which hybridizes under stringent conditions to the full complement of SEQ ID NO:1 ~~or SEQ ID NO:3~~ and is capable of encoding a protein with LysR3 transcriptional regulatory activity, wherein the stringent conditions comprise washing in 5 X SSC at a temperature of ~~from 50 to~~ 68°C; and recovering the produced L-amino acid.

Claim 27 (Currently Amended): The process of Claim 26, wherein said bacterial cell is a *Coryneform bacterium* or *Brevibacterium* ~~*Brevibacterim*~~.

Claim 28 (Currently Amended): The process of Claim 26, wherein said bacterial cell is selected from the group consisting of ~~*Coryneform*~~ *Corynebacterium glutamicum*, *Corynebacterium acetoglutamicum*, *Corynebacterium acetoacidophilum*, *Corynebacterium melassecola*, *Corynebacterium thermoaminogenes*, *Brevibacterium flavum*, *Brevibacterium lactofermentum*, and *Brevibacterium divaricatum*.

Claim 29 (Currently Amended): The process of Claim 26, wherein said bacterial cell is selected from the group consisting of ~~*Coryneform*~~ *Corynebacterium glutamicum* ATCC13032, *Corynebacterium acetoglutamicum* ATCC15806, *Corynebacterium acetoacidophilum* ATCC13870, *Corynebacterium melassecola* ATCC17965,

*Corynebacterium thermoaminogenes* FERM BP-1539, *Brevibacterium flavum* ATCC14067, *Brevibacterium lactofermentum* ATCC13869, and *Brevibacterium divaricatum* ATCC14020.

Claim 30 (Previously Presented) The process of Claim 26, wherein said lysR3 gene prior to being attenuated comprises the polynucleotide sequence of SEQ ID NO:1.

Claim 31 (Currently Amended) The process of Claim 26, wherein said lysR3 gene prior to being attenuated comprises the polynucleotide sequence of SEQ ID NO:3.

Claim 32 (Original): The process of Claim 26, wherein said L-amino acid is L-lysine.

Claim 33 (Cancelled).

Claim 34 (Original) The process of Claim 26, wherein said L-amino acid is L-valine.

Claim 35 (Cancelled).

Claim 36 (Currently Amended): The process of Claim 32, wherein said bacteria further comprises at least one gene ~~whose expression is enhanced~~ which is overexpressed, wherein said gene is selected from the group consisting of dapA, eno, zwf, pyc, and lysE.

Claim 37 (Original): The process of Claim 32, wherein said bacteria further comprises at least one gene whose expression is attenuated, wherein said gene is selected from the group consisting of pck, pgi, and poxB.

Claim 38 (Currently Amended): The process of Claim 26, wherein said bacteria further comprises at least one gene ~~whose expression is enhanced~~ which is overexpressed, wherein said gene is selected from the group consisting of ilvBN, ilvD and mgo.

Claims 39-46 (Cancelled).

Claim 47 (Previously Presented): The process of Claim 26, wherein said lysR3 gene prior to being attenuated comprises a polynucleotide encoding a polypeptide comprising the amino acid sequence of SEQ ID NO:2.